



## INTERNATIONAL APPLICATION PUBLISHED UNDER THE PATENT COOPERATION TREATY (PCT)

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<b>(21) International Application Number:</b> PCT/US99/29976 <b>(22) International Filing Date:</b> 15 December 1999 (15.12.99)  <b>(30) Priority Data:</b> 09/223,586 30 December 1998 (30.12.98) US 09/364,626 29 July 1999 (29.07.99) US  <b>(71) Applicant (for all designated States except US):</b> OLIGOS ETC. INC. [US/US]; 9775 S.W. Commerce Circle, Building C6, Wilsonville, OR 97068 (US).  <b>(72) Inventors; and</b> <b>(75) Inventors/Applicants (for US only):</b> DALE, Roderic, M., K. [US/US]; 26761 S.W. 45th Drive, Wilsonville, OR 97070 (US). ARROW, Amy [US/US]; 184 Birdhill Road, Bethel, ME 04217 (US). THOMPSON, Terry [US/US]; 2222 SW Ek Road, West Linn, OR 97068 (US).  <b>(74) Agent:</b> DEVORE, Dianna, L.; Bozicevic, Field & Francis LLP, 200 Middlefield Road, Suite 200, Menlo Park, CA 94025 (US).		<b>(81) Designated States:</b> AE, AL, AM, AT, AU, AZ, BA, BB, BG, BR, BY, CA, CH, CN, CR, CU, CZ, DE, DK, DM, EE, ES, FI, GB, GD, GE, GH, GM, HR, HU, ID, IL, IN, IS, JP, KE, KG, KP, KR, KZ, LC, LK, LR, LS, LT, LU, LV, MA, MD, MG, MK, MN, MW, MX, NO, NZ, PL, PT, RO, RU, SD, SE, SG, SI, SK, SL, TJ, TM, TR, TT, TZ, UA, UG, US, UZ, VN, YU, ZA, ZW, ARIPO patent (GH, GM, KE, LS, MW, SD, SL, SZ, TZ, UG, ZW), Eurasian patent (AM, AZ, BY, KG, KZ, MD, RU, TJ, TM), European patent (AT, BE, CH, CY, DE, DK, ES, FI, FR, GB, GR, IE, IT, LU, MC, NL, PT, SE), OAPI patent (BF, BJ, CF, CG, CI, CM, GA, GN, GW, ML, MR, NE, SN, TD, TG).  <b>Published</b> <i>Without international search report and to be republished upon receipt of that report.</i>
<b>(54) Title:</b> THERAPEUTIC PHOSPHODIESTERASE INHIBITORS  <b>(57) Abstract</b>  This patent describes the invention of a series of novel therapeutic oligonucleotide polymers targeted at inhibiting expression of genes coding for Phosphodiesterase 4. They are useful as analytical tools in the study of individual PDE isoforms and in the therapeutic treatment of depression, thrombosis, cystic fibrosis, gastric lesions, pulmonary hypertension, glaucoma, multiple sclerosis, atopic dermatitis, asthma and other allergic disorders as well as other illnesses in which an increase of cyclic AMP or a decrease in phosphodiesterase levels is useful.		

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**THERAPEUTIC PHOSPHODIESTERASE INHIBITORS****BACKGROUND OF THE INVENTION**

The enzyme phosphodiesterase (PDE), along with adenylate cyclase and guanylate cyclase, are enzymes responsible for maintaining the correct balance of cyclic AMP and cyclic GMP in cells. There are multiple distinct phosphodiesterases (PDE 1 through PDE 9), most of which exist as two or more isozymes or splice variants that can differ in their cellular distribution, specificity toward hydrolysis of cAMP or cGMP, selective inhibition by various compounds, and sensitivity to regulation by calcium, calmodulin, cAMP, and cGMP (J.A. Beavo in *Cyclic Nucleotide Phosphodiesterases: Structure, Regulation and Drug Action. Multiple Phosphodiesterase Isozymes: Background, Nomenclature, and Implications*. Eds. Beavo, J. and Houslay, M.D., John Wiley and Son, New York, 1990, pp. 3-15 and T.J. Torphy *et al.*, "Novel Phosphodiesterases Inhibitors for the Therapy of Asthma", *Drug News & Prospective*, 6(4) May 1993, pp. 203-214). The PDE4 family, which is specific for cAMP, is composed of at least 4 isozymes (a-d), and multiple splice variants (Houslay, M.D., *et al.* in *Advances in Pharmacology* 44, Eds. J. August *et al.*, p.225, 1998). In total, there may be over 20 PDE4 isoforms expressed in a cell specific pattern regulated by a number of different promoters.

PDE4 is present in the brain and major inflammatory cells and has been found in abnormally elevated levels in a number of diseases including atopic dermatitis or eczema, asthma, and hay fever (*ASTI Connections*, Vol. 8 #1 (1996) p. 3 and *J. of Allergy and Clinical Immunology* 70:452-457,1982). Disease states for which selective PDE4 inhibitors have been sought include: asthma, atopic dermatitis, depression, reperfusion injury, septic shock, toxic shock, endotoxic shock, adult respiratory distress syndrome, autoimmune diabetes, diabetes insipidus, multi-infarct dementia, AIDS, cancer, Crohn's disease, multiple sclerosis, cerebral ischemia, psoriasis, allograft rejection, restenosis, ulcerative colitis, cachexia, cerebral malaria, allergic rhino-conjunctivitis, osteoarthritis, rheumatoid arthritis, chronic bronchitis, eosinophilic granuloma, and autoimmune encephalomyelitis (Houslay *et al.*, 1998). In individuals suffering from atopic diseases, elevated PDE4 activity can be detected in their peripheral blood mononuclear leukocytes, T cells, mast cells, neutrophils and basophils. This increased PDE activity decreases cAMP levels and results in a breakdown of

cAMP control in these cells, which in turn results in increased immune response in the blood and tissues of affected individuals.

PDE inhibitors influence multiple functional pathways, act on multiple immune and inflammatory pathways, and influence synthesis or release of numerous immune mediators

5 (J.M. Hanifin and S.C. Chan, "Atopic Dermatitis-Therapeutic Implication for New Phosphodiesterase Inhibitors, Monocyte Dysregulation of T Cells" in *AACI News*, 7/2, 1995; J.M. Hanifin *et al.*, "Type 4 Phosphodiesterase Inhibitors Have Clinical and In Vitro Anti-inflammatory Effects in Atopic Dermatitis," *J. of Invest. Derm.*, 1996, 107:51-56 and Cohen, V.L. in INC's 7th Annual Conference on Asthma and Allergy (Oct. 27-28, 1997) -

10 *Phosphodiesterase 4 Inhibitors: Second Generation and Beyond*). Clinical use of inhibitors of PDE4 have shown them to be broad spectrum anti-inflammatory agents with impressive activity in models of asthma and other allergic disorders, including atopic dermatitis and hay fever. PDE4 inhibitors that have been used clinically include theophylline, rolipram, denbufylline, CDP 840 (a tri-aryl ethane) and CP80633 (a pyrimidone). PDE4 inhibitors have  
15 been shown to influence eosinophil responses, decrease basophil histamine release, decrease IgE, PGE2, and IL10 synthesis, and decrease anti-CD3 stimulated IL4 production.

All of the PDE4 inhibitors developed to date have been small molecule compounds which exhibit central nervous system and gastrointestinal effects. Although the first generation of PDE4 inhibitors have been effective in reducing PDE4 activity and alleviating a  
20 number of the inflammatory problems caused by overexpression of this enzyme, their effectiveness has been limited by mechanism related side effects and none of them can be used systemically because of nausea and vomiting (*ASTI Connections*, Vol. 8 #1 (1996) p. 3 and Hanifin reference and Cohen, V.L. in INC's 7th Annual Conference on Asthma and Allergy (Oct. 27-28, 1997) - *Phosphodiesterase 4 Inhibitors: Second Generation and Beyond*).

25 Oligonucleotide therapy, i.e., the use of oligonucleotides to modulate the expression of specific genes, offers an opportunity to selectively modify the expression of genes without the undesirable non-specific toxic effects of more traditional therapeutics. The use of antisense oligonucleotides has emerged as a powerful new approach for the treatment of many diseases. The preponderance of the work to date has focused on the use of antisense  
30 oligonucleotides as antiviral agents or as anticancer agents (Wickstrom, E., Ed., *Prospects for Antisense Nucleic Acid Therapy of Cancer and AIDS*, New York: Wiley-Liss, 1991; Crooke, S.T., and Lebleu, B., Eds., *Antisense Research and Applications*, Boca Raton: CRC Press,

1993, pp. 154-182; Baserga, R., and Denhardt, D.T., 1992, *Antisense Strategies*, New York: The New York Academy of Sciences, Vol. 660; Murray, J.A.H., Ed., *Antisense RNA and DNA*, New York: Wiley-Liss, 1993, Agrawal *et al.*, *Proc. Natl. Acad. Sci. USA* 85:7079-7083 (1988), Zamecnik *et al.*, *Proc. Natl. Acad. Sci. USA* 83:4143-4146 (1986)).

5           There is a need to develop a therapeutic agent for modulating PDE4 that is effective but does not possess the side effects of the current PDE4 inhibitors. There is thus a need for a bioavailable, non-toxic therapeutic agent that specifically modulates PDE4.

#### SUMMARY OF THE INVENTION

10           The present invention provides end-blocked acid resistant nucleic acid polymers, *e.g.*, end-blocked 2'-O-alkyl and 2'-O-alkyl-n(O-alkyl) oligonucleotides, for use in modulating PDE4 activity, either *in vivo* or *in vitro*. These PDE4 polymers exhibit substantial stability at low pH, substantial resistance to nuclease degradation, and binding specificity both *in vivo* and *in vitro*. The nucleotides of these polymers may be either ribonucleotides and/or  
15   deoxyribonucleotides, and may be targeted to DNA sequences involved in the expression of PDE4 (*e.g.*, coding sequences, promoter sequences, enhancer sequences, etc.) or to PDE4 mRNA. These low toxicity, highly specific, acid stable, end-blocked polymers represent an improved nucleic acid structure for therapeutic treatments of PDE4-mediated diseases. The 3', 5', or 3' and 5' acid stable, nuclease resistant ends confer improved bioavailability by  
20   increasing nuclease resistance.

          The invention also provides pharmaceutical compositions comprised of polymers of the invention. These pharmaceutical compositions may include any pharmaceutically acceptable carrier. The pharmaceutical composition may also include additives such as adjuvants, stabilizers, fillers and the like.

25           The invention also provides methods of treating a patient in need of such treatment with a therapeutically effective amount of an polymers targeted to PDE4. In a preferred embodiment, the present invention provides a series of antisense oligonucleotide polymers targeted to mRNAs encoding different PDE4 isozymes. The therapeutic effectiveness of an oligonucleotide targeted against PDE4D is demonstrated herein using data from preliminary  
30   human trials.



The invention also comprises methods of reducing the activity of one or more PDE4 enzyme comprising treating a patient with one or more antisense oligonucleotide polymers. Preferably the oligonucleotide is targeted to mRNA coding for PDE4.

5 It is an advantage of the polymers of the invention that the acid stable ends confer an improved stability on the modified nucleic acids in an acidic environment (e.g., the stomach, with a pH of 1 to 2), and thus increase bioavailability of the polymers *in vivo*.

It is another advantage of the polymers of the invention that they bind with specificity to PDE4 target sequences *in vivo* and *in vitro*.

10 It is another advantage of the invention that the end-blocked nucleic acids are non-toxic to a subject treated with the modified nucleic acids. The modified nucleic acids of the present invention, e.g., 2'-O-alkyl and 2'-O-alkyl-n(O-alkyl) end-blocked oligonucleotides, do not display side effects commonly caused by therapeutic administration of regular polyanionic oligonucleotides, such as increased binding to serum and other proteins, stimulation of serum transaminases, decreases in platelet counts, and the like.

15 It is yet another advantage that the acid stable ends confer an improved stability on the PDE4 oligonucleotides in the acid environments of lysosomal vesicles in macrophages and neutrophils.

It is a further advantage that the PDE4 oligonucleotides of the present invention are readily encapsulated in charged liposomes.

20 It is a further advantage that the PDE4 polymers have low toxicity, i.e., mice parenterally treated with a PDE4 oligonucleotide of the invention exhibit an LD<sub>50</sub> of less than one at 400 mg/ml.

25 These and other objects, advantages, and features of the invention will become apparent to those skilled in the art upon reading the details of the nucleic acids and uses thereof as more fully described below.

#### DETAILED DESCRIPTION OF PREFERRED EMBODIMENTS

Before the present invention is described, it is to be understood that this invention is not limited to the particular methodology, protocols, cell lines, animal species or genera,  
30 constructs, and reagents described, as such may, of course, vary. It is also to be understood that the terminology used herein is for the purpose of describing particular embodiments only,

and is not intended to limit the scope of the present invention which will be limited only by the appended claims.

It must be noted that as used herein and in the appended claims, the singular forms "a", "and", and "the" include plural referents unless the context clearly dictates otherwise.

5 Thus, for example, reference to "a construct" includes a plurality of such constructs and reference to "the oligonucleotide" includes reference to one or more oligonucleotides and equivalents thereof known to those skilled in the art, and so forth.

Unless defined otherwise, all technical and scientific terms used herein have the same meaning as commonly understood to one of ordinary skill in the art to which this invention  
10 belongs. Although any methods, devices and materials similar or equivalent to those described herein can be used in the practice or testing of the invention, the preferred methods, devices and materials are now described.

All publications mentioned herein are incorporated herein by reference for the purpose of describing and disclosing, for example, the cell lines, constructs, and methodologies that  
15 are described in the publications which might be used in connection with the presently described invention. The publications discussed above and throughout the text are provided solely for their disclosure prior to the filing date of the present application. Nothing herein is to be construed as an admission that the inventors are not entitled to antedate such disclosure by virtue of prior invention.

20 The publications discussed herein are provided solely for their disclosure prior to the filing date of the present application. Nothing herein is to be construed as an admission that the present invention is not entitled to antedate such publication by virtue of prior invention. Further, the dates of publication provided may be different from the actual publication dates which may need to be independently confirmed.

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## DEFINITIONS

The terms "nucleic acid" and "nucleic acid molecule" as used interchangeably herein, refer to a molecule comprised of nucleotides, i.e., ribonucleotides, deoxyribonucleotides, or both. The term includes monomers and polymers of ribonucleotides and  
30 deoxyribonucleotides, with the ribonucleotide and/or deoxyribonucleotides being connected together, in the case of the polymers, via 5' to 3' linkages. However, linkages may include any of the linkages known in the nucleic acid synthesis art including, for example, nucleic

acids comprising 5' to 2' linkages. The nucleotides used in the nucleic acid molecule may be naturally occurring or may be synthetically produced analogues that are capable of forming base-pair relationships with naturally occurring base pairs. Examples of non-naturally occurring bases that are capable of forming base-pairing relationships include, but are not limited to, aza and deaza pyrimidine analogues, aza and deaza purine analogues, and other heterocyclic base analogues, wherein one or more of the carbon and nitrogen atoms of the purine and pyrimidine rings have been substituted by heteroatoms, e.g., oxygen, sulfur, selenium, phosphorus, and the like.

The term "oligonucleotide" as used herein refers to a nucleic acid molecule comprising from about 1 to about 100 nucleotides, more preferably from 1 to 80 nucleotides, and even more preferably from about 4 to about 35 nucleotides. The term "oligonucleotide polymer" as used herein refers to a polymer of the invention having from about 1 to about 100 nucleotides, more preferably from 1 to 80 nucleotides, and even more preferably from about 4 to about 35 nucleotides.

The term "PDE4 oligonucleotide" as used herein refers to an oligonucleotide that is targeted to sequences that affect PDE4 expression or activity. These include, but are not limited to, PDE4 DNA coding sequences, PDE4 DNA promoter sequences, PDE4 DNA enhancer sequences, mRNA encoding PDE4, and the like.

The terms "modified oligonucleotide" and "modified nucleic acid molecule" as used herein refer to nucleic acids, including oligonucleotides, with one or more chemical modifications at the molecular level of the natural molecular structures of all or any of the nucleic acid bases, sugar moieties, internucleoside phosphate linkages, as well as molecules having added substituents, such as diamines, cholesteryl or other lipophilic groups, or a combination of modifications at these sites. The internucleoside phosphate linkages can be phosphodiester, phosphotriester, phosphoramidate, siloxane, carbonate, carboxymethylester, acetamidate, carbamate, thioether, bridged phosphoramidate, bridged methylene phosphonate, phosphorothioate, methylphosphonate, phosphorodithioate, bridged phosphorothioate and/or sulfone internucleotide linkages, or 3'-3', 2'-5' or 5'-5' linkages, and combinations of such similar linkages (to produce mixed backbone modified oligonucleotides). The modifications can be internal (single or repeated) or at the end(s) of the oligonucleotide molecule and can include additions to the molecule of the internucleoside phosphate linkages, such as cholesteryl, diamine compounds with varying numbers of carbon residues between amino



groups and terminal ribose, deoxyribose and phosphate modifications which cleave or cross-link to the opposite chains or to associated enzymes or other proteins. Electrophilic groups such as ribose-dialdehyde could covalently link with an epsilon amino group of the lysyl-residue of such a protein. A nucleophilic group such as n-ethylmaleimide tethered to an oligomer could covalently attach to the 5' end of an mRNA or to another electrophilic site. The term modified oligonucleotides also includes oligonucleotides comprising modifications to the sugar moieties such as 2'-substituted ribonucleotides, or deoxyribonucleotide monomers, any of which are connected together via 5' to 3' linkages. Modified oligonucleotides may also be comprised of PNA or morpholino modified backbones where target specificity of the sequence is maintained.

The term "nucleic acid backbone" as used herein refers to the structure of the chemical moiety linking nucleotides in a molecule. This may include structures formed from any and all means of chemically linking nucleotides. A modified backbone as used herein includes modifications to the chemical linkage between nucleotides, as well as other modifications that may be used to enhance stability and affinity, such as modifications to the sugar structure. For example an a-anomer of deoxyribose may be used, where the base is inverted with respect to the natural b-anomer. In a preferred embodiment, the 2'-OH of the sugar group may be altered to 2'-O-alkyl or 2'-O-alkyl-n(O-alkyl), which provides resistance to degradation without comprising affinity.

The term "acidification" and "protonation/acidification" as used interchangeably herein refers to the process by which protons (or positive hydrogen ions) are added to proton acceptor sites on a nucleic acid. The proton acceptor sites include the amine groups on the base structures of the nucleic acid and the phosphate of the phosphodiester linkages. As the pH is decreased, the number of these acceptor sites which are protonated increases, resulting in a more highly protonated/acidified nucleic acid.

The term "protonated/acidified nucleic acid" refers to a nucleic acid that, when dissolved in water at a concentration of approximately 16 A<sub>260</sub> per ml, has a pH lower than physiological pH, i.e., lower than approximately pH 7. Modified nucleic acids, nuclease-resistant nucleic acids, and antisense nucleic acids are meant to be encompassed by this definition. Generally, nucleic acids are protonated/acidified by adding protons to the reactive sites on a nucleic acid, although other modifications that will decrease the pH of the nucleic acid can also be used and are intended to be encompassed by this term.

The term "end-blocked" as used herein refers to a nucleic acid with a chemical modification at the molecular level that prevents the degradation of selected nucleotides, e.g., by nuclease action. This chemical modification is positioned such that it protects the integral portion of the nucleic acid, for example the coding region of an antisense oligonucleotide. An end block may be a 3' end block or a 5' end block. For example, a 3' end block may be at the 3'-most position of the molecule, or it may be internal to the 3' ends, provided it is 3' of the integral sequences of the nucleic acid.

The term "substantially nuclease resistant" refers to nucleic acids that are resistant to nuclease degradation, as compared to naturally occurring or unmodified nucleic acids.

Modified nucleic acids of the invention are at least 1.25 times more resistant to nuclease degradation than their unmodified counterpart, more preferably at least 2 times more resistant, even more preferably at least 5 times more resistant, and most preferably at least 10 times more resistant than their unmodified counterpart. Such substantially nuclease resistant nucleic acids include, but are not limited to, nucleic acids with modified backbones such as 2'-halogens, 2'-O-methyl, 2'-O-alkyl, 2'-O-alkyl-n(O-alkyl), 3'-halogens, 3'-O-alkyl, 3'-O-alkyl-n(O-alkyl), phosphodiester linkages, phosphotriester linkages, phosphoramidate linkages, siloxane linkages, carbonate linkages, carboxymethylester linkages, acetamidate linkages, carbamate linkages, thioether linkages, bridged phosphoramidate linkages, bridged methylene phosphonate linkages, phosphorothioate linkages, methylphosphonate linkages, phosphorodithioate linkages, morpholino, bridged phosphorothioate linkages, sulfone internucleotide linkages, 3'-3' linkages, 5'-2' linkages, 5'-5' linkages, 2'-deoxy-erythropentofuranosyl, 2'-fluoro, 2'-O-alkyl nucleotides, 2'-O-alkyl-n(O-alkyl) phosphodiester, morpholino linkages, p-ethoxy oligonucleotides, PNA linkages, p-isopropyl oligonucleotides, or phosphoramidates, sulfide linkages, sulfoxide linkages, sulfamate linkages, sulfamide linkages, formacetal/ ketal linkages, thioformacetal linkages, chiral phosphorous linkages, aminoalkylphosphorothioamidate linkages, and four residue group linkages, or chimeric versions of these backbones.

The term "substantially acid resistant" as used herein refers to nucleic acids that are resistant to acid degradation as compared to unmodified nucleic acids. Typically, the relative acid resistance of a nucleic acid will be measured by comparing the percent degradation of a resistant nucleic acid with the percent degradation of its unmodified counterpart (i.e., a corresponding nucleic acid with "normal" backbone, bases, and phosphodiester linkages). A

nucleic acid that is acid resistant is preferably at least 1.5 times more resistant to acid degradation, at least 2 times more resistant, even more preferably at least 5 times more resistant, and most preferably at least 10 times more resistant than their unmodified counterpart.

5           The term "polymer" as used herein refers to a nucleic acid or oligonucleotide molecule having both 1) a modified nucleic acid backbone and 2) at least one end-block at or near the end of the molecule.

          The term "LD<sub>50</sub>" as used herein is the dose of an active substance that will result in 50 per cent lethality in all treated experimental animals. Although this usually refers to invasive  
10   administration, such as oral, parenteral, and the like, it may also apply to toxicity using less invasive methods of administration, such as topical applications of the active substance.

          The term "alkyl" as used herein refers to a branched or unbranched saturated hydrocarbon chain containing 1-6 carbon atoms, such as methyl, ethyl, propyl, tert-butyl, n-hexyl and the like.

15           The term "sensitivity" as used herein refers to the relative strength of recognition of a nucleic acid by a binding partner, e.g., an oligonucleotide complementary to the nucleic acids of interest, i.e., PDE4. The recognition of the binding partner to the sequence of interest must be significantly greater than the recognition of background sequences, and preferably the strength of recognition of the binding partner is at least 10 times, more preferably at least  
20   100 times greater, and even more preferably at least 500 times greater than recognition of background proteins.

          The term "selectivity" as used herein refers to the preferential binding of a binding partner to a particular nucleic acid sequence. Preferably, a selective binding partner is at least 10 times, more preferably 100 times, and even more preferably 1000 times more likely to bind  
25   to its designated polynucleotide sequence than to any other background sequence.

          The terms "treatment", "treating" and the like are used herein to generally mean obtaining a desired pharmacologic and/or physiologic effect. The effect may be prophylactic in terms of completely or partially preventing a disease or symptom thereof and/or may be therapeutic in terms of a partial or complete cure for a disease and/or adverse effect  
30   attributable to the disease. "Treatment" as used herein covers any treatment of a disease in a mammal, particularly a human, and includes:

(a) preventing a disease from occurring in a subject which may be predisposed to the disease but has not yet been diagnosed as having it;

(b) inhibiting a disease, i.e., arresting its development; or

(c) relieving a disease, i.e., causing regression of the disease. The invention is generally directed toward treating patients by the administration of a nucleic acid sequence that will modulate expression of an endogenous gene *in vivo*.

By "therapeutically effective" amount is meant a nontoxic but sufficient amount of a compound to provide the desired therapeutic effect, in the present case, that dose of modified nucleic acid which will be effective in relieving, ameliorating, or preventing symptoms of the condition or disease being treated.

### GENERAL ASPECTS OF THE INVENTION

The concept of antisense has been well developed and a number of different antisense therapeutic drugs are in or have completed clinical trials. Antisense therapeutic compounds are oligonucleotides, preferably nuclease resistant, complementary to the mRNA coding for a particular protein. Antisense oligonucleotides generally interfere with the transcription or translation of the targeted gene and thereby reduce expression of the target gene. Because of the great specificity that is possible with antisense there are far fewer, if any, side effects.

In a preferred embodiment, the invention uses either neutral or protonated/acidified oligonucleotide polymers that are substantially nuclease resistant. This embodiment includes oligonucleotide polymers completely or partially derivatized by phosphorothioate linkages, 2'-deoxy-erythropentofuranosyl, 2'-fluoro, 2'-O-alkyl or 2'-O-alkyl-n(O-alkyl) phosphodiester, p-ethoxy oligonucleotides, p-isopropyl oligonucleotides, phosphoramidates, chimeric linkages, and any other backbone modifications. This embodiment also includes other modifications that render the oligonucleotide polymers substantially resistant to endogenous nuclease activity. Additional methods of rendering an oligonucleotide polymer nuclease resistant include, but are not limited to, covalently modifying the purine or pyrimidine bases that comprise the oligonucleotide polymer. For example, bases may be methylated, hydroxymethylated, or otherwise substituted (e.g., glycosylated) such that the oligonucleotide polymers comprising the modified bases are rendered substantially nuclease resistant.

In order to enhance the exonuclease resistance of the oligonucleotide polymers of the invention, the 3' and/or 5' ends of the nucleic acid sequence are preferably attached to an



exonuclease blocking function. For example, one or more phosphorothioate nucleotides can be placed at either end of the oligoribonucleotide. Additionally, one or more inverted bases can be placed on either end of the oligonucleotide polymer, or one or more alkyl moieties, e.g., butanol-substituted nucleotides or chemical groups can be placed on one or more ends of the oligonucleotide polymer. 5' and 3' end blocking groups may include one or more phosphorothioate nucleotides (but typically fewer than six), inverted base linkages, or alkyl, alkenyl, or alkynyl groups or substituted nucleotides or 2'-O-alkyl-n(O-alkyl). A partial list of blocking groups includes inverted bases, dideoxynucleotides, methylphosphates, alkyl groups, alcohol groups, aryl groups, cordycepin, cytosine arabanoside, 2'-methoxyethoxy nucleotides, phosphoramidates, a peptide linkage, dinitrophenyl group, 2'- or 3'-O-methyl bases with phosphorothioate linkages, 3'-O-methyl bases, fluorescein, cholesterol, biotin, acridine, rhodamine, psoralen, glyceryl, methyl phosphonates, a hexa-ethyloxy-glycol, butanol, hexanol, and 3'-O-alkyls. An enzyme-resistant butanol preferably has the structure OH-CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub> (4-hydroxybutyl) which is also referred to as a C4 spacer.

Typically, the relative nuclease resistance of a nucleic acid can be measured by comparing the percent digestion of a resistant nucleic acid with the percent digestion of its unmodified counterpart (i.e., a corresponding nucleic acid with "normal" backbone, bases, and phosphodiester linkage). Percent degradation may be determined by using analytical HPLC to assess the loss of full length nucleic acids, or by any other suitable methods (e.g., by visualizing the products on a sequencing gel using staining, autoradiography, fluorescence, etc., or measuring a shift in optical density). Degradation is generally measured as a function of time.

Comparison between unmodified and modified nucleic acids can be made by ratioing the percentage of intact modified nucleic acid to the percentage of intact unmodified nucleic acid. For example, if, after 15 minutes of exposure to a nuclease, 25% (i.e., 75% degraded) of an unmodified nucleic acid is intact, and 50% (i.e., 50% degraded) of a modified nucleic acid is intact, the modified nucleic acid is said to be 2 times (50% divided by 25%) more resistant to nuclease degradation than is the unmodified nucleic acid. Generally, a substantially nuclease resistant nucleic acid will be at least about 1.25 times more resistant to nuclease degradation than an unmodified nucleic acid with a corresponding sequence, typically at least about 1.5 times more resistant, preferably about 2 times more resistant,



more preferably at least about 5 times more resistant, and even more preferably at least about 10 times more resistant after 15 minutes of nuclease exposure.

Percent acid degradation may be determined by using analytical HPLC to assess the loss of full length nucleic acids, or by any other suitable methods (e.g., by visualizing the products on a sequencing gel using staining, autoradiography, fluorescence, etc., or measuring a shift in optical density). Degradation is generally measured as a function of time.

Comparison between unmodified and modified nucleic acids can be made by ratioing the percentage of intact modified nucleic acid to the percentage of intact unmodified nucleic acid. For example, if, after 30 minutes of exposure to a low pH environment, 25% (i.e., 75% degraded) of an unmodified nucleic acid is intact, and 50% (i.e., 50% degraded) of a modified nucleic acid is intact, the modified nucleic acid is said to be 2 times (50% divided by 25%) more resistant to nuclease degradation than is the unmodified nucleic acid. Generally, substantially "acid resistant" nucleic acids will be at least about 1.25 times more resistant to acid degradation than an unmodified nucleic acid with a corresponding sequence, typically at least about 1.5 times more resistant, preferably about 1.75 more resistant, more preferably at least 5 times more resistant and even more preferably at least about 10 times more resistant after 30 minutes of exposure at 37°C to a pH of about 1.5 to about 4.5.

Acidification of nucleic acids is the process by which protons (or hydrogen atoms) are added to the reactive sites on a nucleic acid. As the pH is decreased, the number of reactive sites protonated increases and the result is a more highly protonated/acidified nucleic acid. As the pH of the nucleic acid decreases, its bacterial inhibiting activity increases. Accordingly, the nucleic acids of the invention are protonated/acidified to give a pH when dissolved in water of less than pH 7 to about pH 1, or in preferred embodiments, pH 6 to about 1 or pH 5 to about 1. In other preferred embodiments, the dissolved nucleic acids have a pH from pH 4.5 to about 1 or, in a preferred embodiment, a pH of 4.0 to about 1, or, in a more preferred embodiment, a pH of 3.0 to about 1, or, in another preferred embodiment, a pH of 2.0 to about 1.

A first aspect of the present invention is a method of treating a patient requiring such treatment comprising administering one or more oligonucleotide polymer(s) targeted to hybridize with one or more of the appropriate PDE4 mRNA(s) in a therapeutically effective amount. In a second aspect, the present invention is a method of treating a patient requiring such treatment comprising administering one or more oligonucleotide polymer(s) targeted to

hybridize with one or more of the appropriate PDE4 mRNA(s) in an amount effective to reduce expression of one or more of the PDE4 enzymes. Preferably, the antisense oligonucleotide polymers are in a pharmaceutically acceptable carrier. The targeted PDE4 gene sequence or sequences are selected from the group consisting of the PDE4 genes, their  
5 isozymes and their splice variants.

The methods and compositions of the invention are useful as analytical tools in the study of individual PDE isoforms and in therapeutic treatment. The methods and compositions of the invention are useful for treating various diseases or disorders, including but not limited to: asthma, hay fever, atopic dermatitis, depression, reperfusion injury, septic  
10 shock, toxic shock, endotoxic shock, adult respiratory distress syndrome, autoimmune diabetes, diabetes insipidus, multi-infarct dementia, AIDS, cancer, Crohn's disease, multiple sclerosis, cerebral ischemia, psoriasis, allograft rejection, restenosis, ulcerative colitis, cachexia, cerebral malaria, allergic rhino-conjunctivitis, osteoarthritis, rheumatoid arthritis, chronic bronchitis, eosinophilic granuloma, and autoimmune encephalomyelitis. Particular  
15 embodiments of the present invention are directed towards the treatment of the above diseases.

Optionally, the presently described oligonucleotide polymers may be formulated with a variety of physiological carrier molecules. The presently described oligonucleotide polymers may also be complexed with molecules that enhance their ability to enter the target  
20 cells. Examples of such molecules include, but are not limited to, carbohydrates, polyamines, amino acids, peptides, lipids, and molecules vital to cell growth. For example, the oligonucleotide polymers may be combined with a lipid, the resulting oligonucleotide polymer/lipid emulsion, or liposomal suspension may, *inter alia*, effectively increase the *in vivo* half-life of the oligonucleotide polymer. The use of cationic, anionic, and/or neutral lipid  
25 compositions or liposomes is generally described in International Publications Nos. WO 90/14074, WO 91/16024, WO 91/17424, Pat. No. 4,897,355, herein incorporated by reference.

Alternatively, oligonucleotide polymers directed at PDE targets may also be protonated/acidified to function in a dual role as phosphodiesterase inhibitors and antibacterial  
30 agents. Accordingly, another embodiment of the presently described invention is the use of a PDE modulating therapeutic oligonucleotide polymer that is additionally protonated/acidified

to increase cellular uptake, improve encapsulation in liposomes, so it can also serve as an antibiotic.

Additionally, the oligonucleotide polymer may be complexed with a variety of well established compounds or structures that, for instance, further enhance the *in vivo* stability of the oligonucleotide polymer, or otherwise enhance its pharmacological properties (e.g., increase *in vivo* half-life, reduce toxicity, etc.).

#### Selection of Antisense Oligonucleotide Polymers

Antisense oligonucleotide polymers may be targeted to particular functional domains of a gene or mRNA transcript. The potential targets include, but are not limited to, regulatory regions, the 5'-untranslated region, the translational start site, the translational termination site, the 3'-untranslated region, exon/intron boundaries and splice sites, as well as sequences internal to the coding region.

As a first step in the selection of a target domain within the target gene, the target gene, pre-mRNA, or mRNA sequence is examined for the presence of homopolymer runs (4 or more bases in a row) within the target sequence, the targeting of which should generally be avoided.

After selection of a target gene, marking the homopolymer runs, and identification of a target domain, a region of the pre-mRNA, mRNA or DNA sequence within that domain is analyzed. An analysis of the segment of the sequence extending approximately 5 to 100 bases or more upstream and downstream from a possible antisense target site is performed. The locations of potentially stable secondary structures are defined as stem-loop structures with predicted melting temperatures above 20°C. This analysis can be performed using commercially available software such as OLIGO 4.0 for Mac, or OligoTech. Sequences involved in stem hybridization for loop-stem secondary structure formation are regarded as part of the secondary structure and are treated as a structural unit for the purposes of analysis and selection of an antisense oligonucleotide polymer. In one embodiment of this target selection technique, antisense oligonucleotide polymers can be targeted to sequences of the gene, pre-mRNA, or mRNA with predicted secondary structure melting temperatures of less than 100°C (according to the commercially available analysis programs). In a preferred embodiment, the secondary structure melting temperatures would be less than 60°C. In a further preferred embodiment, the secondary structure melting temperature would be less

than 40°C. In a most preferred embodiment, secondary structure melting temperatures would be less than 20°C.

Oligonucleotide polymer sequences with stable loop structures or which form stable dimers should be avoided. In one embodiment of this target selection technique, antisense oligonucleotide polymers with predicted secondary structure melting temperatures of less than 100°C (according to the commercially available analysis programs) are chosen. In a preferred embodiment the secondary structure melting temperatures would be less than 80°C. In a more preferred embodiment the secondary structure melting temperatures would be less than 60°C. In a further preferred embodiment the secondary structure melting temperatures would be less than 40°C. In the most preferred embodiment, the secondary structure melting temperature is less than 20°C. The formation of stable secondary or tertiary structure by the antisense oligonucleotide polymers may potentially compete with their ability to bind to the target DNA or RNA. Similarly, formation of homodimers by the antisense oligonucleotide polymers may compete with their ability to bind to target DNA or RNA. Homopolymer runs of a single base ideally will be three bases or less within the oligonucleotide polymer sequence. Generally, within a target, those sequences with the lowest secondary structure melting temperatures (loop T<sub>m</sub>), or no secondary structure work best.

Sequences of antisense oligonucleotide polymers useful as compositions and in methods of the present invention include the following:

Target Gene: PDE4A (PDE4A) - - Acc# U68532 (SEQ ID NO:46)

Target Domain  
(Nucleotide  
Position #s)

Antisense Oligonucleotide Sequence

59-35	tta gag cag gtc tcg cag aag aaa t, (SEQ ID NO: 1)
569-545	agc gtc agc atg tat gtc acc atc g, (SEQ ID NO: 2)
886-862	gct tgc tga ggt tct gga aga tgt c, (SEQ ID NO: 3)
1541-1518	aga gct tcc tcg act cct gac aat, (SEQ ID NO: 4)
359-335	atg tta gag ttg ttc agg ctg tta c, (SEQ ID NO: 5)
398-374	agg agc tct tct tga tcg gtc ttc a, (SEQ ID NO: 6)
696-674	gag aat ctc cag gtc cgt gaa ca, (SEQ ID NO: 7)
1350-1333	ggc gct gta gta cca gtc, (SEQ ID NO: 8)

2797-2782           aca ggg aca gag gtc t, (SEQ ID NO: 9)  
 3445-3421           gac ttc tag tca gta tcg cca gga g, (SEQ ID NO: 10)

Target Gene: PDE4B (PDE4B2B) -- Acc# L20971 (SEQ ID NO:47)

5   Target Domain       Antisense Oligonucleotide Sequence  
     (Nucleotide  
     Position #s)

720-696           aca aat cac agt ggt gct ctg cct g, (SEQ ID NO: 11)  
 10   1441-1417       ggg ctt cta aag tca tca tgt agg t, (SEQ ID NO: 12)  
     2185-2166       cta agg tat cga gaa tgt cc, (SEQ ID NO: 13)  
     2506-2482       cat gct cat caa gga tag aat gtt c, (SEQ ID NO: 14)  
     269-247         gac gtt tgg gtt ata taa tac ac, (SEQ ID NO: 15)  
     995-971         ttg tta gaa gcc atc tca ctg aca g, (SEQ ID NO: 16)  
 15   1787-1763       atg tca ata acc atc ttc ctg agt g, (SEQ ID NO: 17)  
     1889-1865       tct agg aga aga agc cct gaa ctt g, (SEQ ID NO: 18)  
     3254-3235       gag ttc tat gca gac tct ca, (SEQ ID NO: 19)  
     3951-3927       tca gta gtt cgg gag cat tca gaa g, (SEQ ID NO: 20)

20   Target Gene: PDE4C (PDE4C1B) -- Acc# Z46632 (SEQ ID NO:48)

Target Domain       Antisense Oligonucleotide Sequence  
 (Nucleotide  
 Position #s)

25   119-102         ttc tcc atg cgc cag aga, (SEQ ID NO: 21)  
     532-515         cag agg agt tcc gag aca, (SEQ ID NO: 22)  
     1681-1664       cga ggc ttg tca cct tct, (SEQ ID NO: 23)  
     2255-2231       tgg ccc taa gtc ctc tgg ttg tcg a, (SEQ ID NO: 24)  
     69-52           gct tgg ctg ctc cta ggc, (SEQ ID NO: 25)  
 30   560-538         atg tcc tct cca tgt agg tcg ct, (SEQ ID NO: 26)  
     1140-1121       gtt ccc act tac gtc cgc ca, (SEQ ID NO: 27)  
     1360-1336       cca agt ctg tga aca cag cct cga g, (SEQ ID NO: 28)  
  
     2003-1979       cga ttg tcc tcc agc gtg tcc agc a, (SEQ ID NO: 29)



3095-3071 agt cat agc tct ctt cag cct cca a, (SEQ ID NO: 30)

Target Gene: PDE4D (PDE4D2A) -- Acc# U50158 (SEQ ID NO:49)

	Target Domain (Nucleotide Position #s)	Antisense Oligonucleotide Sequence
5		
	1683-1659	ttg cac tgt tac gtg tca gga gaa c, (SEQ ID NO: 31)
	1672-1659	cgt gtc agg aga ac, (SEQ ID NO: 32)
10	1645-1621	aca ggc ttc agg ttg gct ttc ctc t, (SEQ ID NO: 33)
	1127-1103	gag gct ttg ttg ggt tgc tca gat c, (SEQ ID NO: 34)
	682-658	aga taa tag cac atg agt aga ctg g, (SEQ ID NO: 35)
	154-130	cat ctc act gac gga gtg cct ggt c, (SEQ ID NO: 36)
	609-592	taa tgg tct tcg aga gtc, (SEQ ID NO: 37)
15	822-798	ttg tac atc aag gca agt tca gag t, (SEQ ID NO: 38)
	1046-1022	gaa gaa ctc cag agc ttg tca ctt t, (SEQ ID NO: 39)
	791-767	tga tca gaa att gat tgg aca cac c, (SEQ ID NO: 40)
	1371-1347	tgg tac cat tca cga ttg tcc tcc a, (SEQ ID NO: 41)

20 Target Gene: PDE4D (PDE4D5A) -- Acc# AF012073 (SEQ ID NO:50)

	Target Domain (Nucleotide Position #s)	Antisense Oligonucleotide Sequence
25	300-276	tct cgg aga gat cac tgg aga gag c, (SEQ ID NO: 42)
	375-351	atg tgc cac cgt gaa acg ccg ctg t, (SEQ ID NO: 43)

Target Gene: PDE4D (PDE4D3) -- Acc# L20970 (SEQ ID NO:51)

	Target Domain (Nucleotide Position #s)	Antisense Oligonucleotide Sequence
30		
	95-76	tcc tac gtt aca tgt agt ga, (SEQ ID NO: 44)
	178-159	cat atc cag gaa tgc ctt ct, (SEQ ID NO: 45)

35

In Vivo Testing of Oligonucleotide Polymers

When used in the therapeutic treatment of disease, an appropriate dosage of an antisense PDE oligonucleotide polymer, or mixture thereof, may be determined by any of several well established methodologies. For instance, animal studies are commonly used to determine the maximal tolerable dose, or MTD, of bio-active agent per kilogram weight. In general, at least one of the animal species tested is mammalian. Those skilled in the art regularly extrapolate doses for efficacy and avoiding toxicity to other species, including human. Additionally, therapeutic dosages may also be altered depending upon factors such as the severity of infection and the size or species of the host.

Oligonucleotide polymers are preferably administered in a pharmaceutically acceptable carrier, via oral, intranasal, rectal, topical, intraperitoneal, intramuscular, subcutaneous, intracranial, subdermal, transdermal, intratracheal methods, or the like.

For example, topical diseases are preferably treated or prevented by formulations designed for topical application. Alternately, where the targeted disease state is in the stomach or gut, preparations of oligonucleotide polymers may be provided by oral dosing. Additionally, pulmonary sites of disease, e.g., asthma, may be treated both parenterally and by direct application of suitably formulated forms of the oligonucleotide polymers to the lung by inhalation therapy.

Where suitably formulated oligonucleotide polymers are administered parenterally, the oligonucleotide polymers can accumulate to relatively high levels in the kidneys, liver, spleen, lymph glands, adrenal gland, aorta, pancreas, bone marrow, heart, and salivary glands. Oligonucleotide polymers also tend to accumulate to a lesser extent in skeletal muscle, bladder, stomach, esophagus, duodenum, fat, and trachea. Still lower concentrations are typically found in the cerebral cortex, brain stem, cerebellum, spinal cord, cartilage, skin, thyroid, and prostate (see generally Crooke, 1993, *Antisense Research and Applications*, CRC Press, Boca Raton, FL).

One of ordinary skill will appreciate that, from a medical practitioner's or patient's perspective, virtually any alleviation or prevention of an undesirable symptom would be desirable. Thus, the terms "treatment", "therapeutic use", or "medicinal use" used herein shall refer to any and all uses of the claimed oligonucleotide polymers that remedy a disease state or symptoms, or otherwise prevent, hinder, retard, or reverse the progression of disease or other undesirable symptoms in any way whatsoever.

Preferably, animal hosts that may be treated using the oligonucleotide polymers of the present invention include, but are not limited to, invertebrates, vertebrates, birds, mammals such as pigs, goats, sheep, cows, dogs, cats, and particularly humans. Oligonucleotide polymers are designed to be appropriate to the particular animal to be treated.

5 Another embodiment of the presently described invention is the use of a therapeutic oligonucleotide polymer that is additionally protonated/acidified so it can also serve as an antibiotic. Purified or crude nucleic acids can be protonated/acidified with acid, including but not limited to, phosphoric acid, nitric acid, hydrochloric acid, acetic acid, etc. For example, acid may be combined with nucleic acids in solution, or alternatively, the nucleic acids may be  
10 dissolved in an acidic solution. Excess acid may be removed by chromatography or in some cases by drying the nucleic acid.

Other procedures to prepare protonated nucleic acids known to the skilled artisan are equally contemplated to be within the scope of the invention. Once the nucleic acids of the present invention have been protonated they may be separated from any undesired  
15 components such as excess acid. The skilled artisan would know of many ways to separate the oligonucleotide polymers from undesired components. For example, the oligonucleotide polymer solution may be subjected to chromatography following protonation. In a preferred embodiment, the oligonucleotide polymer solution is run over a poly(styrene-divinylbenzene) based resin column (e.g., Hamilton's PRP or Polymer Labs' PLRP) following protonation.

20 The protonated/acidified nucleic acids can be used directly, or in a preferred embodiment, processed further to remove any excess acid and salt via precipitation, reverse phase chromatography, diafiltration, or gel filtration. The protonated/acidified oligos can be concentrated by precipitation, lyophilization, solvent evaporation, etc. When suspended in water or saline, the acidified nucleic acid preparations of the invention typically exhibit a pH  
25 between 0.5 and 4.5 depending upon 1) the level of protonation/acidification, which can be determined by how much acid is used in the acidification process, and 2) the concentration of the nucleic acid. Alternatively, nucleic acids can be protonated by passage over a cation exchange column charged with hydrogen ions.

It has been discovered that protonated oligonucleotide polymers, in a sequence  
30 independent manner, exhibit antibacterial properties. The protonated oligonucleotide polymers of the invention are protonated/acidified to give a pH when dissolved in water of less than pH 7 to about pH 1, or in preferred embodiments, pH 6 to about 1 or pH 5 to about

1. In other preferred embodiments, the dissolved nucleic acids have a pH from pH 4.5 to about 1 or, in a preferred embodiment, a pH of 4.0 to about 1, or, in a more preferred embodiment, a pH of 3.0 to about 1, or, in another preferred embodiment, a pH of 2.0 to about 1.

5

#### Pharmaceutical Compositions and Delivery

Oligonucleotide polymers may be formulated with a variety of physiological carrier molecules for *in vivo* use. For example, the oligonucleotide polymers may be combined with a lipid, cationic lipid, or anionic lipid (which may be preferred for protonated/acidified oligonucleotide polymers) and the resulting oligonucleotide polymer/lipid emulsion, or liposomal suspension may, *inter alia*, effectively increase the *in vivo* half-life of the oligonucleotide polymer.

When used in the therapeutic treatment of disease, an appropriate dosage of a PDE4 modulating oligonucleotide polymer, or mixture thereof, may be determined by any of several well established methodologies. For instance, animal studies are commonly used to determine the maximal tolerable dose, or MTD, of bio-active agent per kilogram weight. In general, at least one of the animal species tested is mammalian. Those skilled in the art regularly extrapolate doses for efficacy and avoiding toxicity to other species, including human. Additionally, therapeutic dosages may also be altered depending upon factors such as the severity of the disease, and the size or species of the host.

The use of protonated/acidified oligonucleotide polymers facilitates their encapsulation in anionic lipids which are thought to have a number of advantages over cationic liposomes (R.J. Lee and L. Huang, "Lipidic Vector Systems for Gene Transfer", in *Critical Reviews in Therapeutic Drug Carrier Systems* 14(2):173-206 (1997)). Specifically, anionic liposomes are likely to be less toxic than cationic liposomes, they exhibit lower non-specific uptake, and they can be targeted with the appropriate ligands to specific cells.

Examples of suitable anionic lipids for use with protonated/acidified oligonucleotide polymers include, but are not limited to, cardiolipin, dimyristoyl, dipalmitoyl, or dioleoyl phosphatidyl choline or phosphatidyl glycerol, palmitoyloleoyl phosphatidyl choline or phosphatidyl glycerol, phosphatidic acid, lysophosphatidic acid, phosphatidyl serine, phosphatidyl inositol, and anionic forms of cholesterol. The use of cationic, anionic, and/or neutral lipid compositions or liposomes is generally described in International Publications



Nos. WO 90/14074, WO 91/16024, WO 91/17424, and U.S. Pat. No. 4,897,355, herein incorporated by reference.

By assembling oligonucleotide polymers into lipid-associated structures, the oligonucleotide polymers may be targeted to specific cell types by the incorporation of  
5 suitable targeting agents (i.e., specific antibodies or receptors) into the oligonucleotide polymer/lipid complex.

Pharmaceutical compositions containing oligonucleotide polymers of the invention in intimate admixture with a pharmaceutical carrier can be prepared according to conventional pharmaceutical compounding techniques. The carrier may take a wide variety of forms  
10 depending on the form of preparation desired for administration, e.g., intravenous, oral, topical, aerosol (for topical or inhalation therapy), suppository, parenteral, or spinal injection.

In preparing the compositions in oral dosage form, any of the usual pharmaceutical media may be employed, such as, for example, water, glycols, oils, alcohols, flavoring agents, preservatives, coloring agents and the like in the case of oral liquid preparations (such as, for  
15 example, suspension, elixirs, and solutions); or carriers such as starches, sugars, diluents, granulating agents, lubricants, binders, disintegrating agents and the like in the case of oral solid preparations (such as, for example, powders, capsules and tablets). Because of their ease in administration, tablets and capsules represent the most advantageous oral dosage unit form, in which case solid pharmaceutical carriers are obviously employed. If desired, tablets  
20 may be sugar-coated or enteric-coated by standard techniques.

For parenteral application by injection, preparations may comprise an aqueous solution of a water soluble, or solubilized, and pharmaceutically acceptable form of the oligonucleotide polymer in an appropriate saline solution. Injectable suspensions may also be prepared using appropriate liquid carriers, suspending agents, agents for adjusting the  
25 isotonicity, preserving agents, and the like. Actual methods for preparing parenterally administrable compositions and adjustments necessary for administration to subjects will be shown or apparent to those skilled in the art and are described in more detail in, for example, *Remington's Pharmaceutical Science*, 15th Ed., Mack Publishing Company, Easton, PA (1980), which is incorporated herein by reference. The presently described oligonucleotide  
30 polymers should be parenterally administered at concentrations below the maximal tolerable dose (MTD) established for the oligonucleotides.



For topical administration, the carrier may take a wide variety of forms depending on the preparation, which may be a cream, dressing, gel, lotion, ointment, or liquid.

Aerosols are prepared by dissolving or suspending the oligonucleotide in a propellant such as ethyl alcohol or in propellant and solvent phases. The pharmaceutical compositions for topical or aerosol form will generally contain from about 0.01% by weight (of the oligonucleotide) to about 40% by weight, preferably about (0.02% to about 10% by weight, and more preferably about 0.05% to about 5% by weight depending on the particular form employed.

Suppositories are prepared by mixing the oligonucleotide polymer with a lipid vehicle such as theobroma oil, cacao butter, glycerin, gelatin, or polyoxyethylene glycols.

The presently described oligonucleotide polymers may be administered to the body by virtually any means used to administer conventional therapeutics. A variety of delivery systems are well known in the art for delivering bioactive compounds to an animal. These systems include, but are not limited to, intravenous or intra-muscular or intra-tracheal injection, nasal spray, aerosols for inhalation, and oral or suppository administration. The specific delivery system used depends on the location of the disease, and it is well within the skill of one in the art to determine the location of the disease and to select an appropriate delivery system.

## EXAMPLES

The following examples are put forth so as to provide those of ordinary skill in the art with a complete disclosure and description of how to make and use the subject invention, and are not intended to limit the scope of what is regarded as the invention. Efforts have been made to ensure accuracy with respect to the numbers used (*e.g.*, amounts, temperature, concentrations, *etc.*) but some experimental errors and deviations should be allowed for. Unless otherwise indicated, parts are parts by weight, molecular weight is average molecular weight, temperature is in degrees centigrade, and pressure is at or near ambient.

### EXAMPLE 1: B Cell Mediated Dermatitis

An anti-inflammatory oligonucleotide polymer, OE-2a (SEQ ID NO: 32), is a 2' O-methyl RNA phosphodiester linked, with 5' and 3' ends blocked with inverted Ts, and is targeted against the human PDE4 gene. OE-2a was dissolved at 35 mg/ml (7.7  $\mu$ Molar) in

water at approximately pH 3 and used to treat a 37 year old male with a history of severe recurrent atopic dermatitis. The dermatitis was specifically eradicated with OE-2a. In addition, future occurrences of atopic dermatitis have been completely eliminated.

5 A 37 year old male presented with severe atopic dermatitis, completely covering the inside of both forearms. Atopic dermatitis is a chronic disorder characterized by intensely pruritic inflamed papules. The inflammatory response is associated with over-production of IgE by B lymphocytes. Higher levels of PDE4 have been reported for individuals with atopic dermatitis. An antisense oligonucleotide polymer, OE-2a, specifically targeted to PDE4 was applied to a 2" by 3" segment of the left forearm. A second oligo, OE-1, was applied to  
10 another segment of the same arm. OE-1 has a similar base distribution to OE-2a, but is homologous to a bacterial gene. Oligonucleotide polymers were applied twice at 12 hour intervals.

OE-2a was successful at clearing the dermatitis on the area to which it was applied. OE-1 had no effect. The patient commented prior to the second treatment that the itching  
15 had stopped on the area corresponding to the OE-2a after about 6 hours.

Treatment on the second arm was initiated at this time exclusively with OE-2a. Again there were 2 treatments 12 hours apart. The patient remarked that all itching had ceased within 6 hours. The dermatitis was completely cleared within 12 hours of the second treatment.

20 There have been no recurrences of atopic dermatitis on this individual from February through December of a single year, although he typically suffers from recurring bouts during the heat of the summer.

#### EXAMPLE 2: T Cell Mediated Dermatitis

25 OE-2a at 35 mg/ml (7.7  $\mu$ Molar) in water at approximately pH 3 was used to treat a case of T cell mediated contact dermatitis triggered by poison ivy. The dermatitis was completely eliminated with 2 treatments, and secondary eruptions on other areas of the infected individual were prevented.

A female presented with poison ivy on several areas of the bottom of her foot. The  
30 patient complained of itching that is associated with T cell mediated type hypersensitivity. The contact dermatitis was characterized by redness, induration and vesiculation. The anti-inflammatory oligo OE-2a was applied to the bottom of the foot. Within 12 hours the patient

noted that the intense itching had ceased. A second application of OE-2a resulted in a disappearance of the redness, induration, and vesiculation. It is of interest to note that topical application of OE-2a not only eliminated the visible poison ivy on the bottom of the foot, but also prevented any additional eruptions of poison ivy induced dermatitis elsewhere on the individual.

EXAMPLE 3: Acute Wheal and Flare Reaction

OE-2a at 43.6 mg/ml (7.7  $\mu$ Molar) and approximately pH 7 was able to prevent both an immediate and delayed-type hypersensitivity response when applied within minutes of receiving multiple wasp stings.

A female patient with a history of severe wheal and flare skin reaction in response to insect bites presented with an arm that had received multiple wasp stings. The patient was in severe pain and hysterical. OE-2a was administered immediately. Within 5 minutes the patient was calm and pain free. Remarkably, administration of the OE-2a within 5 minutes of receiving the stings completely prevented any immediate wheal and flare reaction, as well as any delayed reaction.

EXAMPLE 4: Chemically Induced Dermatitis

OE-2a was successfully used to treat a case of chemically induced contact dermatitis that had failed to respond to standard dermatological treatments.

A 59 year old woman presented with a case of contact dermatitis on her neck triggered by a reaction to dry cleaning reagents in a turtle neck sweater. Over the course of a month she had been treated by a dermatologist with cortisone, antibiotics, and a variety of other topical drugs. The various treatments gave no more than temporary sporadic relief and they failed to eliminate the itching and redness. At this time the patient was treated topically with OE-2a. The itching and redness disappeared completely after two treatments with no recurrences over a 5 month period.

EXAMPLE 5: Intranasal Administration of PDE-4

Several healthy male volunteers with severe allergic rhinitis were treated with OE-2a. The males ranged in age from 45 years old to 73 years old. The weights ranged from 175 pounds to 230 pounds. A sterile nasal spray container was filled with a solution containing

OE-2a dissolved in normal saline. The solution of oligonucleotide polymer in saline, at a concentration of concentration of approximately 300 A<sub>260</sub> per ml, was administered via the intranasal route twice daily to each nostril. This dosage regime was continued for three days.

5 Upon examination, following the three day course of treatment, all volunteers were found to be completely free of all symptoms of allergic rhinitis. Specifically there was no nasal congestion, no nasal discharge, no sneezing, no difficulty breathing through the nose, and the nasal passages appeared to be completely clear. There has been no relapse of any of the symptoms of rhinitis since treatment schedule was completed, although the volunteers have continued to be exposed to the allergens that triggered the rhinitis. In conclusion,  
10 intranasal administration of PDE-4 dissolved in saline and administered via an intranasal spray appeared to cure allergic rhinitis in healthy male volunteers after only three days of treatment, with no further recurrences of the symptoms.

While the present invention has been described with reference to the specific  
15 embodiments thereof, it should be understood by those skilled in the art that various changes may be made and equivalents may be substituted without departing from the true spirit and scope of the invention. In addition, many modifications may be made to adapt a particular situation, material, composition of matter, process, process step or steps, to the objective, spirit and scope of the present invention. All such modifications are intended to be within the  
20 scope of the claims appended hereto.

## WHAT IS CLAIMED IS:

1. An acid resistant polymer complementary to PDE4 comprising:  
a nucleic acid having a backbone structure modified from that of a naturally occurring  
5 nucleotide polymer; and  
a blocking chemical modification at or near at least one end of the nucleic acid;  
wherein the polymer is characterized by a pH stability of at least one hour at a pH of  
about 0.01 to about 10 and a nuclease resistance of at least twice that of a naturally occurring  
nucleic acids having the same number of nucleotides.

10 2. The polymer of claim 1, wherein the polymer includes from about two to  
about 100 nucleotides.

3. The polymer of claims 1 or 2, wherein the polymer is targeted to a PDE4  
15 gene selected from the group consisting of: PDE4A1, PDE4A4a, PDE4A4b, PDE4A4c,  
PDE4A5, PDE4A8a, PDE4B1a, PDE4B2a, PDE4B2b, PDE4B2c, PDE4B3a, PDE4C1a,  
PDE4C1b, PDE4C2a, PDE4C3a, PDE4C5a, PDE4C6a, PDE4C7a, PDE4D1a, PDE4D1b,  
PDE4D2a, PDE4D3a, PDE4D3a, PDE4D4a, and PDE4D5a.

20 4. The polymer of claims 1, 2 or 3, wherein the backbone structure includes one  
or more of the group consisting of: wherein the backbone structure includes one or more of  
the group consisting of: 2'-halogens, 2'-O-methyl, 2'-O-alkyl, 2'-O-alkyl-n(O-alkyl), 3'-  
halogens, 3'-O-alkyl, 3'-O-alkyl-n(O-alkyl), phosphodiester linkages, phosphotriester linkages,  
phosphoramidate linkages, siloxane linkages, carbonate linkages, carboxymethylester  
25 linkages, acetamidate linkages, carbamate linkages, thioether linkages, bridged  
phosphoramidate linkages, bridged methylene phosphonate linkages, phosphorothioate  
linkages, methylphosphonate linkages, phosphorodithioate linkages, morpholino, bridged  
phosphorothioate linkages, sulfone internucleotide linkages, 3'-3' linkages, 5'-2' linkages, 5'-5'  
linkages, 2'-deoxy-erythropentofuranosyl, 2'-fluoro, 2'-O-alkyl nucleotides, 2'-O-alkyl-n(O-  
30 alkyl) phosphodiester, morpholino linkages, p-ethoxy oligonucleotides, PNA linkages, p-  
isopropyl oligonucleotides, or phosphoramidates, sulfide linkages, sulfoxide linkages,  
sulfamate linkages, sulfamide linkages, formacetal/ ketal linkages, thioformacetal linkages,



chiral phosphorous linkages, aminoalkylphosphorothioamidate linkages, and four residue group linkages.

5. The polymer of claims 1, 2, 3, or 4, wherein the polymer hybridizes to DNA sequences involved in PDE4 expression.

6. The polymer of claim 1, 2, 3, or 4, wherein the polymer hybridizes to a PDE4 mRNA.

7. The polymer of claims 1, 3 or 4, wherein the nucleotides have a sequence selected from the group consisting of: SEQ ID NO: 1; SEQ ID NO: 2; SEQ ID NO: 3; SEQ ID NO: 4; SEQ ID NO: 5; SEQ ID NO: 6; SEQ ID NO: 7; SEQ ID NO: 8; SEQ ID NO: 9; SEQ ID NO: 10; SEQ ID NO: 11; SEQ ID NO: 12; SEQ ID NO: 13; SEQ ID NO: 14; SEQ ID NO: 15; SEQ ID NO: 16; SEQ ID NO: 17; SEQ ID NO: 18; SEQ ID NO: 19; SEQ ID NO: 20; SEQ ID NO: 21; SEQ ID NO: 22; SEQ ID NO: 23; SEQ ID NO: 24; SEQ ID NO: 25; SEQ ID NO: 26; SEQ ID NO: 27; SEQ ID NO: 28; SEQ ID NO: 29; SEQ ID NO: 30; SEQ ID NO: 31; SEQ ID NO: 32; SEQ ID NO: 33; SEQ ID NO: 34; SEQ ID NO: 35; SEQ ID NO: 36; SEQ ID NO: 37; SEQ ID NO: 38; SEQ ID NO: 39; SEQ ID NO: 40; SEQ ID NO: 41; SEQ ID NO: 42; SEQ ID NO: 43; SEQ ID NO: 44; and SEQ ID NO: 45.

8. The polymer of claims 1, 2, 3, 4, 5, 6 or 7, wherein the polymer is protonated to have a pH at or below 7.

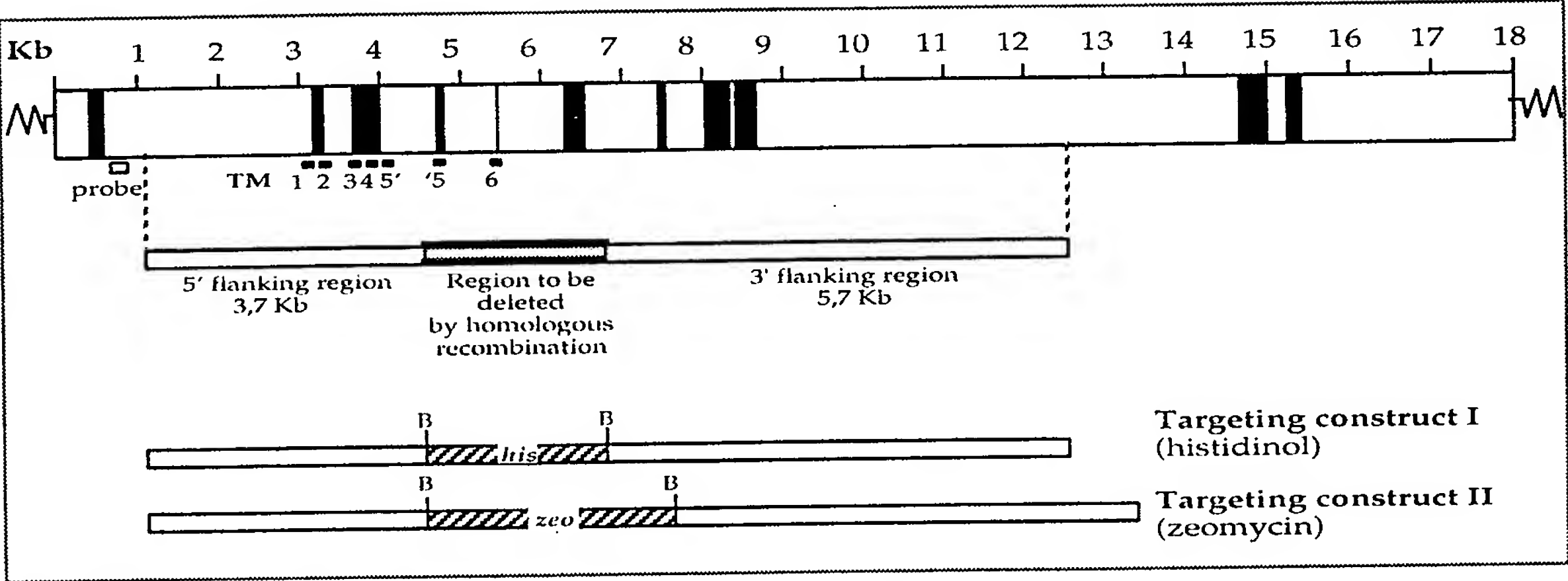
9. A pharmaceutical composition comprised of a polymer of nucleotides directed to modulating PDE4, said polymer having a nucleic acid backbone structure modified from that of a naturally occurring nucleic acid and a blocking chemical modification at or near at least one end of the polymer; and

a pharmaceutically acceptable carrier.

10. The pharmaceutical composition of claim 9, wherein said polymer includes from about one to about 100 nucleotides.

11. The pharmaceutical composition of claims 9 or 10, wherein said polymer is protonated to have a pH at or below 7.

FIGURE 1.



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<110> Dale, Roderic M. K.  
Arrow, Amy  
Thompson, Terry

<120> Antisense Phosphodiesterase Inhibitors

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Asp	Asn	Tyr	Ser	Asp	Arg	Ile	Gln	Val	Leu	Gln	Asn	Leu	Val	His	Cys
	530					535					540				
Ala	Asp	Leu	Ser	Asn	Pro	Thr	Lys	Pro	Leu	Pro	Leu	Tyr	Arg	Gln	Trp
545				550						555					560
Thr	Asp	Arg	Ile	Met	Ala	Glu	Phe	Phe	Gln	Gln	Gly	Asp	Arg	Glu	Arg
			565						570					575	
Glu	Ser	Gly	Leu	Asp	Ile	Ser	Pro	Met	Cys	Asp	Lys	His	Thr	Ala	Ser
			580					585					590		
Val	Glu	Lys	Ser	Gln	Val	Gly	Phe	Ile	Asp	Tyr	Ile	Ala	His	Pro	Leu
	595						600					605			
Trp	Glu	Thr	Trp	Ala	Asp	Leu	Val	His	Pro	Asp	Ala	Gln	Asp	Leu	Leu
	610					615					620				
Asp	Thr	Leu	Glu	Asp	Asn	Arg	Glu	Trp	Tyr	Gln	Ser	Lys	Ile	Pro	Arg
625					630					635					640
Ser	Pro	Ser	Asp	Leu	Thr	Asn	Pro	Glu	Arg	Asp	Gly	Pro	Asp	Arg	Phe
				645					650					655	

Gln Phe Glu Leu Thr Leu Glu Glu Ala Glu Glu Glu Asp Glu Glu Glu  
 660 665 670  
 Glu Glu Glu Gly Glu Glu Thr Ala Leu Ala Lys Glu Ala Leu Glu Leu  
 675 680 685  
 Pro Asp Thr Glu Leu Leu Ser Pro Glu Ala Gly Pro Asp Pro Gly Asp  
 690 695 700  
 Leu Pro Leu Asp Asn Gln Arg Thr  
 705 710

<210> 48  
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 <212> PRT  
 <213> human

<400> 48

Met Lys Glu His Gly Gly Thr Phe Ser Ser Thr Gly Ile Ser Gly Gly  
 1 5 10 15  
 Ser Gly Asp Ser Ala Met Asp Ser Leu Gln Pro Leu Gln Pro Asn Tyr  
 20 25 30  
 Met Pro Val Cys Leu Phe Ala Glu Glu Ser Tyr Gln Lys Leu Ala Met  
 35 40 45  
 Glu Thr Leu Glu Glu Leu Asp Trp Cys Leu Asp Gln Leu Glu Thr Ile  
 50 55 60  
 Gln Thr Tyr Arg Ser Val Ser Glu Met Ala Ser Asn Lys Phe Lys Arg  
 65 70 75 80  
 Met Leu Asn Arg Glu Leu Thr His Leu Ser Glu Met Ser Arg Ser Gly  
 85 90 95  
 Asn Gln Val Ser Glu Tyr Ile Ser Asn Thr Phe Leu Asp Lys Gln Asn  
 100 105 110  
 Asp Val Glu Ile Pro Ser Pro Thr Gln Lys Asp Arg Glu Lys Lys Lys  
 115 120 125  
 Lys Gln Gln Leu Met Thr Gln Ile Ser Gly Val Lys Lys Leu Met His  
 130 135 140  
 Ser Ser Ser Leu Asn Asn Thr Ser Ile Ser Arg Phe Gly Val Asn Thr  
 145 150 155 160  
 Glu Asn Glu Asp His Leu Ala Lys Glu Leu Glu Asp Leu Asn Lys Trp  
 165 170 175  
 Gly Leu Asn Ile Phe Asn Val Ala Gly Tyr Ser His Asn Arg Pro Leu  
 180 185 190  
 Thr Cys Ile Met Tyr Ala Ile Phe Gln Glu Arg Asp Leu Leu Lys Thr  
 195 200 205  
 Phe Arg Ile Ser Ser Asp Thr Phe Ile Thr Tyr Met Met Thr Leu Glu  
 210 215 220  
 Asp His Tyr His Ser Asp Val Ala Tyr His Asn Ser Leu His Ala Ala  
 225 230 235 240  
 Asp Val Ala Gln Ser Thr His Val Leu Leu Ser Thr Pro Ala Leu Asp  
 245 250 255  
 Ala Val Phe Thr Asp Leu Glu Ile Leu Ala Ala Ile Phe Ala Ala Ala  
 260 265 270  
 Ile His Asp Val Asp His Pro Gly Val Ser Asn Gln Phe Leu Ile Asn  
 275 280 285  
 Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn Asp Glu Ser Val Leu Glu  
 290 295 300  
 Asn His His Leu Ala Val Gly Phe Lys Leu Leu Gln Glu Glu His Cys  
 305 310 315 320  
 Asp Ile Phe Met Asn Leu Thr Lys Lys Gln Arg Gln Thr Leu Arg Lys  
 325 330 335  
 Met Val Ile Asp Met Val Leu Ala Thr Asp Met Ser Lys His Met Ser  
 340 345 350  
 Leu Leu Ala Asp Leu Lys Thr Met Val Glu Thr Lys Lys Val Thr Ser  
 355 360 365  
 Ser Gly Val Leu Leu Leu Asp Asn Tyr Thr Asp Arg Ile Gln Val Leu  
 370 375 380

Arg Asn Met Val His Cys Ala Asp Leu Ser Asn Pro Thr Lys Ser Leu  
 385 390 395 400  
 Glu Leu Tyr Arg Gln Trp Thr Asp Arg Ile Met Glu Glu Phe Phe Gln  
 405 410 415  
 Gln Gly Asp Lys Glu Arg Glu Arg Gly Met Glu Ile Ser Pro Met Cys  
 420 425 430  
 Asp Lys His Thr Ala Ser Val Glu Lys Ser Gln Val Gly Phe Ile Asp  
 435 440 445  
 Tyr Ile Val His Pro Leu Trp Glu Thr Trp Ala Asp Leu Val Gln Pro  
 450 455 460  
 Asp Ala Gln Asp Ile Leu Asp Thr Leu Glu Asp Asn Arg Asn Trp Tyr  
 465 470 475 480  
 Gln Ser Met Ile Pro Gln Ser Pro Ser Pro Pro Leu Asp Glu Gln Asn  
 485 490 495  
 Arg Asp Cys Gln Gly Leu Met Glu Lys Phe Gln Phe Glu Leu Thr Leu  
 500 505 510  
 Asp Glu Glu Asp Ser Glu Gly Pro Glu Lys Glu Gly Glu Gly His Ser  
 515 520 525  
 Tyr Phe Ser Ser Thr Lys Thr Leu Cys Val Ile Asp Pro Glu Asn Arg  
 530 535 540  
 Asp Ser Leu Gly Glu Thr Asp Ile Asp Ile Ala Thr Glu Asp Lys Ser  
 545 550 555 560  
 Pro Val Asp Thr

<210> 49  
 <211> 507  
 <212> PRT  
 <213> human

<400> 49  
 Met Ala Ser Asn Lys Phe Lys Arg Met Leu Asn Arg Glu Leu Thr His  
 1 5 10 15  
 Leu Ser Glu Met Ser Arg Ser Gly Asn Gln Val Ser Glu Phe Ile Ser  
 20 25 30  
 Asn Thr Phe Leu Asp Lys Gln His Glu Val Glu Ile Pro Ser Pro Thr  
 35 40 45  
 Gln Lys Glu Lys Glu Lys Lys Lys Arg Pro Met Ser Gln Ile Ser Gly  
 50 55 60  
 Val Lys Lys Leu Met His Ser Ser Ser Leu Thr Asn Ser Ser Ile Pro  
 65 70 75 80  
 Arg Phe Gly Val Lys Thr Glu Gln Glu Asp Val Leu Ala Lys Glu Leu  
 85 90 95  
 Glu Asp Val Asn Lys Trp Gly Leu His Val Phe Arg Ile Ala Glu Leu  
 100 105 110  
 Ser Gly Asn Arg Pro Leu Thr Val Ile Met His Thr Ile Phe Gln Glu  
 115 120 125  
 Arg Asp Leu Leu Lys Thr Phe Lys Ile Pro Val Asp Thr Leu Ile Thr  
 130 135 140  
 Tyr Leu Met Thr Leu Glu Asp His Tyr His Ala Asp Val Ala Tyr His  
 145 150 155 160  
 Asn Asn Ile His Ala Ala Asp Val Val Gln Ser Thr His Val Leu Leu  
 165 170 175  
 Ser Thr Pro Ala Leu Glu Ala Val Phe Thr Asp Leu Glu Ile Leu Ala  
 180 185 190  
 Ala Ile Phe Ala Ser Ala Ile His Asp Val Asp His Pro Gly Val Ser  
 195 200 205  
 Asn Gln Phe Leu Ile Asn Thr Asn Ser Glu Leu Ala Leu Met Tyr Asn  
 210 215 220  
 Asp Ser Ser Val Leu Glu Asn His His Leu Ala Val Gly Phe Lys Leu  
 225 230 235 240  
 Leu Gln Glu Glu Asn Cys Asp Ile Phe Gln Asn Leu Thr Lys Lys Gln  
 245 250 255



Arg	Gln	Ser	Leu	Arg	Lys	Met	Val	Ile	Asp	Ile	Val	Leu	Ala	Thr	Asp
			260					265					270		
Met	Ser	Lys	His	Met	Asn	Leu	Leu	Ala	Asp	Leu	Lys	Thr	Met	Val	Glu
		275					280					285			
Thr	Lys	Lys	Val	Thr	Ser	Ser	Gly	Val	Leu	Leu	Leu	Asp	Asn	Tyr	Ser
	290					295					300				
Asp	Arg	Ile	Gln	Val	Leu	Gln	Asn	Met	Val	His	Cys	Ala	Asp	Leu	Ser
305					310					315					320
Asn	Pro	Thr	Lys	Pro	Leu	Gln	Leu	Tyr	Arg	Gln	Trp	Thr	Asp	Arg	Ile
			325						330					335	
Met	Glu	Glu	Phe	Phe	Arg	Gln	Gly	Asp	Arg	Glu	Arg	Glu	Arg	Gly	Met
			340					345					350		
Glu	Ile	Ser	Pro	Met	Cys	Asp	Lys	His	Asn	Ala	Ser	Val	Glu	Lys	Ser
		355					360					365			
Gln	Val	Gly	Phe	Ile	Asp	Tyr	Ile	Val	His	Pro	Leu	Trp	Glu	Thr	Trp
	370				375						380				
Ala	Asp	Leu	Val	His	Pro	Asp	Ala	Gln	Asp	Ile	Leu	Asp	Thr	Leu	Glu
385					390				395						400
Asp	Asn	Arg	Glu	Trp	Tyr	Gln	Ser	Thr	Ile	Pro	Gln	Ser	Pro	Ser	Pro
			405						410					415	
Ala	Pro	Asp	Asp	Pro	Glu	Glu	Gly	Arg	Gln	Gly	Gln	Thr	Glu	Lys	Phe
			420					425					430		
Gln	Phe	Glu	Leu	Thr	Leu	Glu	Glu	Asp	Gly	Glu	Ser	Asp	Thr	Glu	Lys
		435					440					445			
Asp	Ser	Gly	Ser	Gln	Val	Glu	Glu	Asp	Thr	Ser	Cys	Ser	Asp	Ser	Lys
	450					455					460				
Thr	Leu	Arg	Thr	Gln	Asp	Ser	Glu	Ser	Thr	Glu	Ile	Pro	Leu	Asp	Glu
465					470					475					480
Gln	Val	Glu	Glu	Glu	Ala	Val	Gly	Glu	Glu	Glu	Glu	Ser	Gln	Pro	Glu
			485					490						495	
Ala	Cys	Val	Ile	Asp	Asp	Arg	Ser	Pro	Asp	Thr					
			500					505							

<210> 50  
 <211> 745  
 <212> PRT  
 <213> human

<400> 50

Met	Ala	Gln	Gln	Thr	Ser	Pro	Asp	Thr	Leu	Thr	Val	Pro	Glu	Val	Asp
1				5					10					15	
Asn	Pro	His	Cys	Pro	Asn	Pro	Trp	Leu	Asn	Glu	Asp	Leu	Val	Lys	Ser
			20					25					30		
Leu	Arg	Glu	Asn	Leu	Leu	Gln	His	Glu	Lys	Ser	Lys	Thr	Ala	Arg	Lys
		35					40					45			
Ser	Val	Ser	Pro	Lys	Leu	Ser	Pro	Val	Ile	Ser	Pro	Arg	Asn	Ser	Pro
	50					55					60				
Arg	Leu	Leu	Arg	Arg	Met	Leu	Leu	Ser	Ser	Asn	Ile	Pro	Lys	Gln	Arg
65					70					75					80
Arg	Phe	Thr	Val	Ala	His	Thr	Cys	Phe	Asp	Val	Asp	Asn	Gly	Thr	Ser
				85					90					95	
Ala	Gly	Arg	Ser	Pro	Leu	Asp	Pro	Met	Thr	Ser	Pro	Gly	Ser	Gly	Leu
			100					105					110		
Ile	Leu	Gln	Ala	Asn	Phe	Val	His	Ser	Gln	Arg	Arg	Glu	Ser	Phe	Leu
		115					120					125			
Tyr	Arg	Ser	Asp	Ser	Asp	Tyr	Asp	Leu	Ser	Pro	Lys	Ser	Met	Ser	Arg
	130				135						140				
Asn	Ser	Ser	Ile	Ala	Ser	Asp	Ile	His	Gly	Asp	Asp	Leu	Ile	Val	Thr
145					150					155					160
Pro	Phe	Ala	Gln	Val	Leu	Ala	Ser	Leu	Arg	Thr	Val	Arg	Asn	Asn	Phe
			165					170						175	
Ala	Ala	Leu	Thr	Asn	Leu	Gln	Asp	Arg	Ala	Pro	Ser	Lys	Arg	Ser	Pro
			180					185					190		

Met	Cys	Asn	Gln	Pro	Ser	Ile	Asn	Lys	Ala	Thr	Ile	Thr	Glu	Glu	Ala
		195					200					205			
Tyr	Gln	Lys	Leu	Ala	Ser	Glu	Thr	Leu	Glu	Glu	Leu	Asp	Trp	Cys	Leu
	210					215					220				
Asp	Gln	Leu	Glu	Thr	Leu	Gln	Thr	Arg	His	Ser	Val	Ser	Glu	Met	Ala
225					230					235					240
Ser	Asn	Lys	Phe	Lys	Arg	Met	Leu	Asn	Arg	Glu	Leu	Thr	His	Leu	Ser
				245					250					255	
Glu	Met	Ser	Arg	Ser	Gly	Asn	Gln	Val	Ser	Glu	Phe	Ile	Ser	Asn	Thr
			260					265					270		
Phe	Leu	Asp	Lys	Gln	His	Glu	Val	Glu	Ile	Pro	Ser	Pro	Thr	Gln	Lys
		275						280				285			
Glu	Lys	Glu	Lys	Lys	Lys	Arg	Pro	Met	Ser	Gln	Ile	Ser	Gly	Val	Lys
	290					295					300				
Lys	Leu	Met	His	Ser	Ser	Ser	Leu	Thr	Asn	Ser	Ser	Ile	Pro	Arg	Phe
305					310					315					320
Gly	Val	Lys	Thr	Glu	Gln	Glu	Asp	Val	Leu	Ala	Lys	Glu	Leu	Glu	Asp
				325					330					335	
Val	Asn	Lys	Trp	Gly	Leu	His	Val	Phe	Arg	Ile	Ala	Glu	Leu	Ser	Gly
			340					345					350		
Asn	Arg	Pro	Leu	Thr	Val	Ile	Met	His	Thr	Ile	Phe	Gln	Glu	Arg	Asp
		355					360					365			
Leu	Leu	Lys	Thr	Phe	Lys	Ile	Pro	Val	Asp	Thr	Leu	Ile	Thr	Tyr	Leu
	370					375					380				
Met	Thr	Leu	Glu	Asp	His	Tyr	His	Ala	Asp	Val	Ala	Tyr	His	Asn	Asn
385					390					395					400
Ile	His	Ala	Ala	Asp	Val	Val	Gln	Ser	Thr	His	Val	Leu	Leu	Ser	Thr
				405					410					415	
Pro	Ala	Leu	Glu	Ala	Val	Phe	Thr	Asp	Leu	Glu	Ile	Leu	Ala	Ala	Ile
			420					425					430		
Phe	Ala	Ser	Ala	Ile	His	Asp	Val	Asp	His	Pro	Gly	Val	Ser	Asn	Gln
		435					440					445			
Phe	Leu	Ile	Asn	Thr	Asn	Ser	Glu	Leu	Ala	Leu	Met	Tyr	Asn	Asp	Ser
	450					455					460				
Ser	Val	Leu	Glu	Asn	His	His	Leu	Ala	Val	Gly	Phe	Lys	Leu	Leu	Gln
465					470					475					480
Glu	Glu	Asn	Cys	Asp	Ile	Phe	Gln	Asn	Leu	Thr	Lys	Lys	Gln	Arg	Gln
				485				490						495	
Ser	Leu	Arg	Lys	Met	Val	Ile	Asp	Ile	Val	Leu	Ala	Thr	Asp	Met	Ser
			500					505					510		
Lys	His	Met	Asn	Leu	Leu	Ala	Asp	Leu	Lys	Thr	Met	Val	Glu	Thr	Lys
		515					520					525			
Lys	Val	Thr	Ser	Ser	Gly	Val	Leu	Leu	Leu	Asp	Asn	Tyr	Ser	Asp	Arg
	530					535					540				
Ile	Gln	Val	Leu	Gln	Asn	Met	Val	His	Cys	Ala	Asp	Leu	Ser	Asn	Pro
545					550					555					560
Thr	Lys	Pro	Leu	Gln	Leu	Tyr	Arg	Gln	Trp	Thr	Asp	Arg	Ile	Met	Glu
				565					570					575	
Glu	Phe	Phe	Arg	Gln	Gly	Asp	Arg	Glu	Arg	Glu	Arg	Gly	Met	Glu	Ile
			580					585					590		
Ser	Pro	Met	Cys	Asp	Lys	His	Asn	Ala	Ser	Val	Glu	Lys	Ser	Gln	Val
		595					600					605			
Gly	Phe	Ile	Asp	Tyr	Ile	Val	His	Pro	Leu	Trp	Glu	Thr	Trp	Ala	Asp
	610					615					620				
Leu	Val	His	Pro	Asp	Ala	Gln	Asp	Ile	Leu	Asp	Thr	Leu	Glu	Asp	Asn
625					630					635					640
Arg	Glu	Trp	Tyr	Gln	Ser	Thr	Ile	Pro	Gln	Ser	Pro	Ser	Pro	Ala	Pro
				645					650					655	
Asp	Asp	Pro	Glu	Glu	Gly	Arg	Gln	Gly	Gln	Thr	Glu	Lys	Phe	Gln	Phe
			660					665					670		
Glu	Leu	Thr	Leu	Glu	Glu	Asp	Gly	Glu	Ser	Asp	Thr	Glu	Lys	Asp	Ser
		675					680					685			

Gly	Ser	Gln	Val	Glu	Glu	Asp	Thr	Ser	Cys	Ser	Asp	Ser	Lys	Thr	Leu
690						695					700				
Cys	Thr	Gln	Asp	Ser	Glu	Ser	Thr	Glu	Ile	Pro	Leu	Asp	Glu	Gln	Val
705					710					715					720
Glu	Glu	Glu	Ala	Val	Gly	Glu	Glu	Glu	Glu	Ser	Gln	Pro	Glu	Ala	Cys
				725					730						735
Val	Ile	Asp	Asp	Arg	Ser	Pro	Asp	Thr							
			740					745							

<210> 51  
 <211> 673  
 <212> PRT  
 <213> human

<400> 51

Met	Met	His	Val	Asn	Asn	Phe	Pro	Phe	Arg	Arg	His	Ser	Trp	Ile	Cys
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Phe	Asp	Val	Asp	Asn	Gly	Thr	Ser	Ala	Gly	Arg	Ser	Pro	Leu	Asp	Pro
			20					25					30		
Met	Thr	Ser	Pro	Gly	Ser	Gly	Leu	Ile	Leu	Gln	Ala	Asn	Phe	Val	His
		35					40					45			
Ser	Gln	Arg	Arg	Glu	Ser	Phe	Leu	Tyr	Arg	Ser	Asp	Ser	Asp	Tyr	Asp
	50					55					60				
Leu	Ser	Pro	Lys	Ser	Met	Ser	Arg	Asn	Ser	Ser	Ile	Ala	Ser	Asp	Ile
65				70					75						80
His	Gly	Asp	Asp	Leu	Ile	Val	Thr	Pro	Phe	Ala	Gln	Val	Leu	Ala	Ser
				85					90						95
Leu	Arg	Thr	Val	Arg	Asn	Asn	Phe	Ala	Ala	Leu	Thr	Asn	Leu	Gln	Asp
			100					105							110
Arg	Ala	Pro	Ser	Lys	Arg	Ser	Pro	Met	Cys	Asn	Gln	Pro	Ser	Ile	Asn
		115					120					125			
Lys	Ala	Thr	Ile	Thr	Glu	Glu	Ala	Tyr	Gln	Lys	Leu	Ala	Ser	Glu	Thr
	130					135					140				
Leu	Glu	Glu	Leu	Asp	Trp	Cys	Leu	Asp	Gln	Leu	Glu	Thr	Leu	Gln	Thr
145				150					155						160
Arg	His	Ser	Val	Ser	Glu	Met	Ala	Ser	Asn	Lys	Phe	Lys	Arg	Met	Leu
				165					170					175	
Asn	Arg	Glu	Leu	Thr	His	Leu	Ser	Glu	Met	Ser	Arg	Ser	Gly	Asn	Gln
			180					185					190		
Val	Ser	Glu	Phe	Ile	Ser	Asn	Thr	Phe	Leu	Asp	Lys	Gln	His	Glu	Val
		195					200					205			
Glu	Ile	Pro	Ser	Pro	Thr	Gln	Lys	Glu	Lys	Glu	Lys	Lys	Lys	Arg	Pro
	210					215					220				
Met	Ser	Gln	Ile	Ser	Gly	Val	Lys	Lys	Leu	Met	His	Ser	Ser	Ser	Leu
225				230					235						240
Thr	Asn	Ser	Ser	Ile	Pro	Arg	Phe	Gly	Val	Lys	Thr	Glu	Gln	Glu	Asp
				245					250						255
Val	Leu	Ala	Lys	Glu	Leu	Glu	Asp	Val	Asn	Lys	Trp	Gly	Leu	His	Val
			260					265					270		
Phe	Arg	Ile	Ala	Glu	Leu	Ser	Gly	Asn	Arg	Pro	Leu	Thr	Val	Ile	Met
		275					280					285			
His	Thr	Ile	Phe	Gln	Glu	Arg	Asp	Leu	Leu	Lys	Thr	Phe	Lys	Ile	Pro
	290					295					300				
Val	Asp	Thr	Leu	Ile	Thr	Tyr	Leu	Met	Thr	Leu	Glu	Asp	His	Tyr	His
305					310					315					320
Ala	Asp	Val	Ala	Tyr	His	Asn	Asn	Ile	His	Ala	Ala	Asp	Val	Val	Gln
				325					330						335
Ser	Thr	His	Val	Leu	Leu	Ser	Thr	Pro	Ala	Leu	Glu	Ala	Val	Phe	Thr
			340					345					350		
Asp	Leu	Glu	Ile	Leu	Ala	Ala	Ile	Phe	Ala	Ser	Ala	Ile	His	Asp	Val
		355					360					365			
Asp	His	Pro	Gly	Val	Ser	Asn	Gln	Phe	Leu	Ile	Asn	Thr	Asn	Ser	Glu
	370					375						380			

Leu	Ala	Leu	Met	Tyr	Asn	Asp	Ser	Ser	Val	Leu	Glu	Asn	His	His	Leu
385					390					395					400
Ala	Val	Gly	Phe	Lys	Leu	Leu	Gln	Glu	Glu	Asn	Cys	Asp	Ile	Phe	Gln
				405				410						415	
Asn	Leu	Thr	Lys	Lys	Gln	Arg	Gln	Ser	Leu	Arg	Lys	Met	Val	Ile	Asp
			420					425					430		
Ile	Val	Leu	Ala	Thr	Asp	Met	Ser	Lys	His	Met	Asn	Leu	Leu	Ala	Asp
		435				440					445				
Leu	Lys	Thr	Met	Val	Glu	Thr	Lys	Lys	Val	Thr	Ser	Ser	Gly	Val	Leu
	450					455					460				
Leu	Leu	Asp	Asn	Tyr	Ser	Asp	Arg	Ile	Gln	Val	Leu	Gln	Asn	Met	Val
465					470					475					480
His	Cys	Ala	Asp	Leu	Ser	Asn	Pro	Thr	Lys	Pro	Leu	Gln	Leu	Tyr	Arg
				485					490					495	
Gln	Trp	Thr	Asp	Arg	Ile	Met	Glu	Glu	Phe	Phe	Arg	Gln	Gly	Asp	Arg
			500				505						510		
Glu	Arg	Glu	Arg	Gly	Met	Glu	Ile	Ser	Pro	Met	Cys	Asp	Lys	His	Asn
		515				520						525			
Ala	Ser	Val	Glu	Lys	Ser	Gln	Val	Gly	Phe	Ile	Asp	Tyr	Ile	Val	His
	530					535					540				
Pro	Leu	Trp	Glu	Thr	Trp	Ala	Asp	Leu	Val	His	Pro	Asp	Ala	Gln	Asp
545					550					555					560
Ile	Leu	Asp	Thr	Leu	Glu	Asp	Asn	Arg	Glu	Trp	Tyr	Gln	Ser	Thr	Ile
			565						570					575	
Pro	Gln	Ser	Pro	Ser	Pro	Ala	Pro	Asp	Asp	Pro	Glu	Glu	Gly	Arg	Gln
			580					585					590		
Gly	Gln	Thr	Glu	Lys	Phe	Gln	Phe	Glu	Leu	Thr	Leu	Glu	Glu	Asp	Gly
		595				600						605			
Glu	Ser	Asp	Thr	Glu	Lys	Asp	Ser	Gly	Ser	Gln	Val	Glu	Glu	Asp	Thr
	610					615					620				
Ser	Cys	Ser	Asp	Ser	Lys	Thr	Leu	Cys	Thr	Gln	Asp	Ser	Glu	Ser	Thr
625					630					635					640
Glu	Ile	Pro	Leu	Asp	Glu	Gln	Val	Glu	Glu	Glu	Ala	Val	Gly	Glu	Glu
			645					650						655	
Glu	Glu	Ser	Gln	Pro	Glu	Ala	Cys	Val	Ile	Asp	Asp	Arg	Ser	Pro	Asp
			660					665					670		
Thr															